

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 101734,661C  
Source: TEW16  
Date Processed by STIC: 4/27/07

***ENTERED***



IFW16

RAW SEQUENCE LISTING DATE: 04/27/2007  
 PATENT APPLICATION: US/10/734,661C TIME: 11:40:55

Input Set : A:\81408-4400 sequence listing.txt  
 Output Set: N:\CRF4\04272007\J734661C.raw

3 <110> APPLICANT: Yayon, Avner  
 4 Rom, Eran  
 5 Thomassen-Wolf, Elisabeth  
 6 Borges, Eric  
 8 <120> TITLE OF INVENTION: ANTIBODIES THAT BLOCK RECEPTOR PROTEIN TYROSINE  
 KINASE ACTIVATION,  
 9 . METHODS OF SCREENING AND USES THEREOF  
 11 <130> FILE REFERENCE: 81408-4400  
 13 <140> CURRENT APPLICATION NUMBER: US 10/734,661C (pg. 6)  
 14 <141> CURRENT FILING DATE: 2003-12-15  
 16 <150> PRIOR APPLICATION NUMBER: US 60/299,187  
 17 <151> PRIOR FILING DATE: 2001-06-20  
 19 <150> PRIOR APPLICATION NUMBER: PCT/IL02/00494  
 20 <151> PRIOR FILING DATE: 2002-06-20  
 22 <160> NUMBER OF SEQ ID NOS: 106  
 24 <170> SOFTWARE: PatentIn version 3.2  
 26 <210> SEQ ID NO: 1  
 27 <211> LENGTH: 806  
 28 <212> TYPE: PRT  
 29 <213> ORGANISM: Homo sapiens  
 31 <300> PUBLICATION INFORMATION:  
 32 <308> DATABASE ACCESSION NO: np\_000133  
 33 <309> DATABASE ENTRY DATE: 2001-02-21  
 34 <313> RELEVANT RESIDUES: (1)..(806)  
 36 <400> SEQUENCE: 1  
 38 Met Gly Ala Pro Ala Cys Ala Leu Ala Leu Cys Val Ala Val Ala Ile  
 39 1 5 10 15  
 42 Val Ala Gly Ala Ser Ser Glu Ser Leu Gly Thr Glu Gln Arg Val Val  
 43 20 25 30  
 46 Gly Arg Ala Ala Glu Val Pro Gly Pro Glu Pro Gly Gln Gln Glu Gln  
 47 35 40 45  
 50 Leu Val Phe Gly Ser Gly Asp Ala Val Glu Leu Ser Cys Pro Pro Pro  
 51 50 55 60  
 54 Gly Gly Gly Pro Met Gly Pro Thr Val Trp Val Lys Asp Gly Thr Gly  
 55 65 70 75 80  
 58 Leu Val Pro Ser Glu Arg Val Leu Val Gly Pro Gln Arg Leu Gln Val  
 59 85 90 95  
 62 Leu Asn Ala Ser His Glu Asp Ser Gly Ala Tyr Ser Cys Arg Gln Arg  
 63 100 105 110  
 66 Leu Thr Gln Arg Val Leu Cys His Phe Ser Val Arg Val Thr Asp Ala  
 67 115 120 125  
 70 Pro Ser Ser Gly Asp Asp Glu Asp Gly Glu Asp Glu Ala Glu Asp Thr  
 71 130 135 140  
 74 Gly Val Asp Thr Gly Ala Pro Tyr Trp Thr Arg Pro Glu Arg Met Asp

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75	145	150	155	160
78	Lys Lys Leu Leu Ala Val Pro Ala Ala Asn Thr Val Arg Phe Arg Cys			
79		165	170	175
82	Pro Ala Ala Gly Asn Pro Thr Pro Ser Ile Ser Trp Leu Lys Asn Gly			
83		180	185	190
86	Arg Glu Phe Arg Gly Glu His Arg Ile Gly Gly Ile Lys Leu Arg His			
87		195	200	205
90	Gln Gln Trp Ser Leu Val Met Glu Ser Val Val Pro Ser Asp Arg Gly			
91		210	215	220
94	Asn Tyr Thr Cys Val Val Glu Asn Lys Phe Gly Ser Ile Arg Gln Thr			
95		225	230	235
98	Tyr Thr Leu Asp Val Leu Glu Arg Ser Pro His Arg Pro Ile Leu Gln			240
99		245	250	255
102	Ala Gly Leu Pro Ala Asn Gln Thr Ala Val Leu Gly Ser Asp Val Glu			
103		260	265	270
106	Phe His Cys Lys Val Tyr Ser Asp Ala Gln Pro His Ile Gln Trp Leu			
107		275	280	285
110	Lys His Val Glu Val Asn Gly Ser Lys Val Gly Pro Asp Gly Thr Pro			
111		290	295	300
114	Tyr Val Thr Val Leu Lys Thr Ala Gly Ala Asn Thr Thr Asp Lys Glu			
115		305	310	315
118	Leu Glu Val Leu Ser Leu His Asn Val Thr Phe Glu Asp Ala Gly Glu			320
119		325	330	335
122	Tyr Thr Cys Leu Ala Gly Asn Ser Ile Gly Phe Ser His His Ser Ala			
123		340	345	350
126	Trp Leu Val Val Leu Pro Ala Glu Glu Glu Leu Val Glu Ala Asp Glu			
127		355	360	365
130	Ala Gly Ser Val Tyr Ala Gly Ile Leu Ser Tyr Gly Val Gly Phe Phe			
131		370	375	380
134	Leu Phe Ile Leu Val Val Ala Ala Val Thr Leu Cys Arg Leu Arg Ser			
135		385	390	395
138	Pro Pro Lys Lys Gly Leu Gly Ser Pro Thr Val His Lys Ile Ser Arg			400
139		405	410	415
142	Phe Pro Leu Lys Arg Gln Val Ser Leu Glu Ser Asn Ala Ser Met Ser			
143		420	425	430
146	Ser Asn Thr Pro Leu Val Arg Ile Ala Arg Leu Ser Ser Gly Glu Gly			
147		435	440	445
150	Pro Thr Leu Ala Asn Val Ser Glu Leu Glu Leu Pro Ala Asp Pro Lys			
151		450	455	460
154	Trp Glu Leu Ser Arg Ala Arg Leu Thr Leu Gly Lys Pro Leu Gly Glu			
155		465	470	475
158	Gly Cys Phe Gly Gln Val Val Met Ala Glu Ala Ile Gly Ile Asp Lys			480
159		485	490	495
162	Asp Arg Ala Ala Lys Pro Val Thr Val Ala Val Lys Met Leu Lys Asp			
163		500	505	510
166	Asp Ala Thr Asp Lys Asp Leu Ser Asp Leu Val Ser Glu Met Glu Met			
167		515	520	525
170	Met Lys Met Ile Gly Lys His Lys Asn Ile Ile Asn Leu Leu Gly Ala			
171		530	535	540

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174 Cys Thr Gln Gly Gly Pro Leu Tyr Val Leu Val Glu Tyr Ala Ala Lys  
 175 545 550 555 560  
 178 Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg Arg Pro Pro Gly Leu Asp  
 179 565 570 575  
 182 Tyr Ser Phe Asp Thr Cys Lys Pro Pro Glu Glu Gln Leu Thr Phe Lys  
 183 580 585 590  
 186 Asp Leu Val Ser Cys Ala Tyr Gln Val Ala Arg Gly Met Glu Tyr Leu  
 187 595 600 605  
 190 Ala Ser Gln Lys Cys Ile His Arg Asp Leu Ala Ala Arg Asn Val Leu  
 191 610 615 620  
 194 Val Thr Glu Asp Asn Val Met Lys Ile Ala Asp Phe Gly Leu Ala Arg  
 195 625 630 635 640  
 198 Asp Val His Asn Leu Asp Tyr Tyr Lys Lys Thr Thr Asn Gly Arg Leu  
 199 645 650 655  
 202 Pro Val Lys Trp Met Ala Pro Glu Ala Leu Phe Asp Arg Val Tyr Thr  
 203 660 665 670  
 206 His Gln Ser Asp Val Trp Ser Phe Gly Val Leu Leu Trp Glu Ile Phe  
 207 675 680 685  
 210 Thr Leu Gly Gly Ser Pro Tyr Pro Gly Ile Pro Val Glu Glu Leu Phe  
 211 690 695 700  
 214 Lys Leu Leu Lys Glu Gly His Arg Met Asp Lys Pro Ala Asn Cys Thr  
 215 705 710 715 720  
 218 His Asp Leu Tyr Met Ile Met Arg Glu Cys Trp His Ala Ala Pro Ser  
 219 725 730 735  
 222 Gln Arg Pro Thr Phe Lys Gln Leu Val Glu Asp Leu Asp Arg Val Leu  
 223 740 745 750  
 226 Thr Val Thr Ser Thr Asp Glu Tyr Leu Asp Leu Ser Ala Pro Phe Glu  
 227 755 760 765  
 230 Gln Tyr Ser Pro Gly Gly Gln Asp Thr Pro Ser Ser Ser Ser Gly  
 231 770 775 780  
 234 Asp Asp Ser Val Phe Ala His Asp Leu Leu Pro Pro Ala Pro Pro Ser  
 235 785 790 795 800  
 238 Ser Gly Gly Ser Arg Thr  
 239 805  
 242 <210> SEQ ID NO: 2  
 243 <211> LENGTH: 32  
 244 <212> TYPE: DNA  
 245 <213> ORGANISM: Artificial Sequence  
 247 <220> FEATURE:  
 248 <223> OTHER INFORMATION: artificial primer  
 250 <400> SEQUENCE: 2  
 251 acgtgctagc tgagtccttg gggacggagc ag 32  
 254 <210> SEQ ID NO: 3  
 255 <211> LENGTH: 55  
 256 <212> TYPE: DNA  
 257 <213> ORGANISM: Artificial Sequence  
 259 <220> FEATURE:  
 260 <223> OTHER INFORMATION: artificial primer  
 262 <400> SEQUENCE: 3

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263 acgtctcgag ttaatggta tggtgatggt gtgcatacac acagcccgcc tcgtc  
 266 <210> SEQ ID NO: 4  
 267 <211> LENGTH: 1147  
 268 <212> TYPE: DNA  
 269 <213> ORGANISM: Homo sapiens  
 271 <300> PUBLICATION INFORMATION:  
 272 <308> DATABASE ACCESSION NO: m58051  
 273 <309> DATABASE ENTRY DATE: 1994-11-08  
 274 <313> RELEVANT RESIDUES: (1)..(1147)  
 276 <400> SEQUENCE: 4  
 277 ggcgcgtgcc tgaggacgcc gcggcccccg ccccgccat gggcgccct gcctgcgccc 60  
 279 tcgcgtctg cgtggccgtg gccatcggtt ccggcgccct ctcggagtcc ttggggacgg 120  
 281 agcagcgcgt cgtggggcga gcggcagaag tccccggccc agagcccgcc cagcaggagc 180  
 283 agttggtctt cggcagcggg gatgtgtt agctgagctg tccccggccc ggggtggtc 240  
 285 ccatggggcc cactgtctgg gtcaaggatg gcacagggtt ggtgcctcg gagcgtgtcc 300  
 287 tggggggcc ccagcggctg cagggtgtga atgcctccca cgaggactcc ggggcctaca 360  
 289 gctgccggca ggggtcacg cagcgcgtac tggccactt cagtgtgcgg gtgacagacg 420  
 291 ctccatcctc gggagatgac gaagacgggg aggacgaggc tgaggacaca ggtgtggaca 480  
 293 cagggggccc ttactggaca cggcccgagc ggtggacaa gaagctgtg gccgtggccg 540  
 295 ccgccaacac cgtccgttc cgctggccag ccgctggcaa cccactccc tccatctct 600  
 297 ggtgaagaa cggcagggag ttccggggcg agcaccgcat tggaggcata aagctgcggc 660  
 299 atacgcgtg gggctggc atggaaagcg tggccctcg ggaccggcc aactacacct 720  
 301 gctcgtgga gaacaagttt ggcagcatcc ggcagacgta cacgctgac gtgtggagc 780  
 303 gctcccccgcga cggcccatc ctgcaggccg ggctggccgc caaccagacg ggggtgtgg 840  
 305 gcagcgacgt ggagtccac tgcaagggtt acagtgcgc acagccccc atccagtggc 900  
 307 tcaaggacgt ggaggtaaac ggcagcaagg tggggccgga cggcacaccc tacgttaccg 960  
 309 tgctcaagac ggcggcgct aacaccaccc acaaggagct agagggttctc tccttgcaca 1020  
 311 acgtcacctt tgaggacgcc ggggagataca cctgcctggc gggcaattct attgggttt 1080  
 313 ctcatcactc tgcgtggctg gtgggtgtgc cagccgagga ggagctggg gaggctgacg 1140  
 315 aggccggg  
 318 <210> SEQ ID NO: 5  
 319 <211> LENGTH: 5695  
 320 <212> TYPE: DNA  
 321 <213> ORGANISM: EXPRESSION VECTOR pCEP-PU/ACT7 *Invalid Response* ↙ ↘  
 323 <400> SEQUENCE: 5  
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 326 ccgcatagtt aagccaggat ctgctccctg cttgtgtt ggaggtcgct gagtagtgcg 120  
 328 cgagcaaaat ttaagctaca acaaggcaag gcttgcggca caattgcatt aagaatctgc 180  
 330 tttaggttag gctttgcg ctgcttcgcg atgtacgggc cagatatacg cgttgacatt 240  
 332 gattattgac tagttattaa tagtaatcaa ttacgggtc attagttcat agccatata 300  
 334 tggagttccg cgttacataa cttacggtaa atggccgc tggctgaccg cccaaacgacc 360  
 336 cccgcccatt gacgtcaata atgacgtatg ttcccatagt aacgccaata gggactttcc 420  
 338 attgacgtca atgggtggac tatttacggtaa aactgccc cttggcagta catcaagtgt 480  
 340 atcatatgcc aagtacggcc cctattgacg tcaatgacgg taaatggccc gcctggcatt 540  
 342 atgcccagta catgaccta tggacttcc tctacttgcgtt gatcatctac gtattagtca 600  
 344 tcgttattac catggtgatg cgggtttggc agtacatcaa tggcgtgga tagcggtttg 660  
 346 actcacgggg atttccaagt ctccacccca ttgacgtcaa tggagttt gtttggcacc 720  
 348 aaaatcaacg ggacttcca aatgtcgta acaactccgc cccattgacg caaatggcg 780  
 350 gtaggcgtgt acgggtggag gtctatataa qcaqagctct ctqqctaact aqqaaccca 840

Invalid Response

← P15  
see item  
60  
120  
180  
240  
300  
360  
420  
480  
540  
600  
#10 on  
error  
summary  
sheet

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/734,661C

DATE: 04/27/2007

TIME: 11:40:55

Input Set : A:\81408-4400 sequence listing.txt

Output Set: N:\CRF4\04272007\J734661C.raw

352	ctgcttactg	gcttatacgaa	attaatacga	ctcactatacg	ggagacccaa	gctggctagc	900
354	gtttaaactt	aagcttggta	ccgagctcgg	atccccgtcg	tgcatactatc	gaaggtcg	960
356	gagatcccgaa	ggagccaaa	tcttgtgaca	aaactcacac	atgcccaccc	tgcccagcac	1020
358	ctgaactcct	ggggggaccg	tcagtcttcc	tcttcccccc	aaaacccaaag	gacaccctca	1080
360	tgtatctcccg	gaccctcgag	gtcacatcg	tgggtgtgga	cgtgagccac	gaagaccctg	1140
362	aggtaaagt	caactggta	gtggacggcg	tggaggtgca	taatgccaag	acaaggccgc	1200
364	gggaggagca	gtacaacagc	acgtaccggg	tggtcagcg	cctcaccgtc	ctgcaccagg	1260
366	actggctgaa	tggcaaggag	tacaagtgc	aggtctccaa	caaagccctc	ccagccccc	1320
368	tcgagaaaac	catctccaa	gccaaagggc	agcccccaga	accacagggt	tacaccctgc	1380
370	cccatcccg	ggatgagctg	accaagaacc	aggtcagcct	gacctgcctg	gtcaaaggct	1440
372	tctatcccg	cgacatcgcc	gtggagtggg	agagcaatgg	gcagccggag	aacaactaca	1500
374	agaccacgccc	tcccgtgctg	gactccgacg	gctccttctt	cctctacagc	aagctcaccg	1560
376	tggacaagag	caggtggcag	caggggaacg	tcttctcatg	ctccgtgatg	catgaggctc	1620
378	tgcacaacca	ctacacgcag	aagagcctct	ccctgtctcc	gggttaatatg	tctagagggc	1680
380	ccgtttaaac	ccgctgatca	gcctcgactg	tgccttctag	ttgcccagcca	tctttgttt	1740
382	gcccctcccc	cgtgccttcc	ttgaccctgg	aaggtgccac	tcccactgtc	ctttccta	1800
384	aaaatgagga	aattgcatcg	cattgtctga	gtaggtgtca	ttctattctg	gggggtgggg	1860
386	tggggcagga	cagcaagggg	gaggattggg	aagacaatag	caggcatgt	ggggatgcgg	1920
388	tggctctat	ggcttctgag	gccccaaagaa	ccagctgggg	ctctaggggg	tatccccacg	1980
390	cgcctgttag	cggcgattt	agcgcggcg	gtgtgtggg	tacgcgcagc	gtgaccgcta	2040
392	cacttgcag	cgccttagcg	cccgctcctt	tcgcttctt	cccttcctt	ctcgccacgt	2100
394	tcgcccgtt	tcccgtaa	gctctaaatc	ggggcatccc	tttagggttc	cgatttagt	2160
396	ctttacggca	cctcgacccc	aaaaaaacttg	attagggtga	ttgttcacgt	agtggccat	2220
398	cgcctgtata	gacggttttt	cgcctttga	cgttggagtc	cacgttctt	aatagtggac	2280
400	tcttgttcca	aactggaaaca	acactcaacc	ctatctcggt	ctattcttt	gattataag	2340
402	ggattttggg	gatttggcc	tattgtttaa	aaaatgagct	gatttaacaa	aaatttaacg	2400
404	cgaattaatt	ctgtggatg	tgtgtcagtt	agggtgtgga	aagtccccag	gctccccagg	2460
406	caggcagaag	tatgcaaaagc	atgcatactca	attagtcagc	aaccagggt	ggaaagtccc	2520
408	caggctcccc	agcaggcaga	atgatgcaaa	gcatgcac	caattagtca	gcaaccatag	2580
410	tcccgcccc	aactccgccc	atccgcccc	taactccgccc	cagttccgccc	cattctccgc	2640
412	cccatggctg	actaattttt	tttatttatg	cagaggccga	ggccgcctt	gcctctgagc	2700
414	tattccagaa	gtagtggagga	ggctttttt	gaggcctagg	ctttgtcaaa	aagctcccg	2760
416	gagcttgtat	atccattttc	ggatctgatc	agcacgtgtt	gacaattat	catggcata	2820
418	gtatatcggc	atagtataat	acgacaagg	gaggaactaa	accatggcca	agttgaccag	2880
420	tgccgttccg	gtgctcaccg	cgcgcgacgt	cgcggagcg	gtcgagttt	ggaccgaccg	2940
422	gctcgggtt	tcccggtact	tctgtggagga	cgacttcgc	ggtgtggtcc	gggacgacgt	3000
424	gaccctgtt	atcagcgccg	tccaggacca	ggtggcccg	gacaacaccc	tggcttgggt	3060
426	gtgggtgcgc	ggcctggacg	agctgtacgc	cgagtggcg	gaggtcg	ccacacaactt	3120
428	ccgggacgcc	tccggccgg	ccatgaccga	gatcgccgag	cagccgtggg	ggcgggagtt	3180
430	cgcctgtgc	gaccggccg	gcaactgcgt	gcacttcgt	gccgaggagc	aggactgaca	3240
432	cgtgctacga	gatttcgatt	ccaccgcgc	tttctatgaa	aggttgggt	tcggaatcg	3300
434	tttccgggac	gcccgttgg	tgtatcttca	gcccggggat	ctcatgtcg	agtttctcg	3360
436	ccaccccaac	ttgtttattt	cagcttataa	tggttacaaa	taaagcaata	gcatcacaaa	3420
438	tttcacaaat	aaagcatttt	tttctactgca	ttcttagttt	ggtttgtcca	aactcatcaa	3480
440	tgtatcttat	catgtctgt	taccgtcgac	ctctagctag	agcttggcg	aatcatggc	3540
442	atagctgtt	cctgtgtgaa	attgttatcc	gctcacaattt	ccacacaaca	tacgagccgg	3600
444	aagcataaaag	tgtaaaggct	gggggtgccta	atgagtgagc	taactcacat	taattgcgtt	3660
446	gcgcgtactg	cccgcttcc	agtccggaaa	cctgtcg	cagctgcatt	aatgaatcg	3720
448	ccaaacgcgc	gggagaggcg	gtttgcgtat	tggcgctt	tccgcttct	cgctcactga	3780

RAW SEQUENCE LISTING ERROR SUMMARY                   DATE: 04/27/2007  
PATENT APPLICATION: US/10/734,661C                   TIME: 11:40:56

Input Set : A:\81408-4400 sequence listing.txt  
Output Set: N:\CRF4\04272007\J734661C.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:54; N Pos. 253,254,255  
Seq#:56; N Pos. 256,257,258  
Seq#:70; N Pos. 1,2,3  
Seq#:74; N Pos. 1,2,3  
Seq#:81; N Pos. 1,2,3  
Seq#:83; N Pos. 1,2,3

**VERIFICATION SUMMARY** DATE: 04/27/2007  
PATENT APPLICATION: US/10/734,661C TIME: 11:40:56

Input Set : A:\81408-4400 sequence listing.txt  
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L:1612 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54 after pos.:240  
L:1662 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56 after pos.:240  
L:1968 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 after pos.:0  
L:2064 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74 after pos.:0  
L:2234 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 after pos.:0  
L:2286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83 after pos.:0